

TNF & IFN-g 223 genes up-regulated in Y1 cells 16 hr

Fold	Cont	TNF+IFNg	Gene name	Acc#
2.2	50	109	Acid beta glucosidase {IMAGE:331646}	W08086.1
2.1	76	157	AMP deaminase 3 {IMAGE:750480}	AA389897.1
2.1	53	109	Arachidonate 12-lipoxygenase, pseudogene 2 {IMAGE:335369}	W36511.1
2.4	115	281	B-cell leukemia/lymphoma 3 {IMAGE:716976}	AA266002.1
2.1	83	172	B-cell translocation gene 2, anti-proliferative {IMAGE:583186}	AA154848.1
2	53	105	Cathelin-like protein {IMAGE:334755}	W17771.1
2	40	79	CCAAT/enhancer binding protein alpha (C/EBP), related sequence 1 {IM	AA473938.1
2	54	106	Coagulation factor II {IMAGE:350344}	W40725.1
2.2	40	88	Crystallin, gamma S {IMAGE:483498}	AI893668.1
2.2	43	94	Cytochrome P450, 2b13, phenobarbital inducible, type c {IMAGE:316281}	AI325330.1
2.1	68	141	EST {IMAGE:474319}	AA038807.1
2.6	409	1083	ESTs {IMAGE:315676}	W09957.1
2.4	434	1047	ESTs {IMAGE:316914}	W11926.1
2.1	51	106	ESTs {IMAGE:317515}	W34080.1
2.1	151	316	ESTs {IMAGE:318969}	W34482.1
2.2	37	82	ESTs {IMAGE:330233}	W14737.1
2.3	66	154	ESTs {IMAGE:334410}	W18278.1
2.7	245	663	ESTs {IMAGE:350182}	W34722.1
2.1	53	111	ESTs {IMAGE:372368}	W53544.1
2	258	517	ESTs {IMAGE:373112}	W62546.1
2	65	133	ESTs {IMAGE:385744}	W64241.1
2.1	65	136	ESTs {IMAGE:403413}	W82905.1
2.9	59	170	ESTs {IMAGE:404589}	W82894.1
2	544	1090	ESTs {IMAGE:419146}	W88005.1
2.1	101	217	ESTs {IMAGE:427480}	AA002452.1
2	324	663	ESTs {IMAGE:436999}	AA002783.1
2.3	69	159	ESTs {IMAGE:459562}	AA027381.1
2.1	61	131	ESTs {IMAGE:466371}	AA033504.1

ESTs {IMAGE:467873}
2.4 121 285
2 54 109
2.1 60 124
2.7 291 772
2.1 77 158
2 69 135
3.2 64 204
2.2 95 208
6.2 38 234
4.6 194 896
9.7 50 484
2.5 41 102
4.9 26 128
2.1 46 95
2.1 161 341
17.1 22 376
3.6 36 129
2 50 99
2.1 82 171
3.7 266 995
3.4 209
3.4 41 138
2.2 53 118
2.3 68 156
17.1 161 341
3.6 36 129
2 50 99
2.1 82 171
3.7 266 995
3.4 209
3.4 41 138
2.2 53 118
2.3 68 156
2.1 49 102
2 45 88
2 61 122
2 53 107
2.3 104 238
3.4 45 153
4 393 1563

AA036517.1
AI322694.1
AA049610.1
AA061982.1
AI552474.1
AI595201.1
AA107718.1
AA138025.1
AA143986.1
AA140542.1
AA145136.1
AA154419.1
AA168416.1
AA174434.1
AA174941.1
AA172624.1
AA189578.1
AA198524.1
AA230444.1
AA250462.1
AA396152.1
AA388221.1
AA387356.1
AA386832.1
AA387164.1
AA387199.1
AA415579.1
AA415691.1
AA472833.1
AA473234.1
AA473205.1
AA466508.1

2.4	61	ESTs {IMAGE:818790}	AA467382.1
	2	ESTs {IMAGE:831741}	AA444941.1
2.3	38	ESTs {IMAGE:833477}	AA465972.1
2.6	62	ESTs {IMAGE:834617}	AA467546.1
2.1	48	ESTs {IMAGE:835536}	AA422515.1
2.1	43	ESTs {IMAGE:836133}	AA426897.1
3.5	59	ESTs {IMAGE:837565}	AA432818.1
2	47	ESTs {IMAGE:845839}	AA416091.1
2	80	ESTs {IMAGE:872986}	AA466235.1
2.1	99	ESTs {IMAGE:876106}	AA475528.1
2	50	ESTs {IMAGE:904836}	AA522117.1
3.6	118	ESTs {IMAGE:945014}	AA537234.1
2.8	48	ESTs {IMAGE:961094}	AA547202.1
2.1	147	ESTs, Highly similar to CYSTATHIONINE GAMMA-LYASE [Homo sapier	AB892342.1
2.5	59	ESTs, Highly similar to CYSTATHIONINE GAMMA-LYASE [Homo sapier	AA245993.1
5.7	55	ESTs, Highly similar to HYPOTHETICAL 103.5 KD PROTEIN IN RAD26-{	AI639807.1
4	580	ESTs, Highly similar to INTERFERON-INDUCIBLE PROTEIN [Rattus nor	AA048837.1
6.3	40	ESTs, Highly similar to LYSYL-TRNA SYNTHETASE [Cricetulus longica	AA175815.1
4.1	165	ESTs, Highly similar to NUCLEAR FACTOR NF-KAPPA-B P100 SUBUNIT	AI385509.1
3	149	ESTs, Highly similar to SERINE HYDROXYMETHYLTRANSFERASE, MIT	AA208877.1
2.2	152	ESTs, Highly similar to UTP--GLUCOSE-1-PHOSPHATE URIDYLYLTRAI	AA473123.1
2.3	389	ESTs, Highly similar to 54 kDa oligoadenylylate synthetase-like protein p:	AA123007.1
2.3	49	ESTs, Highly similar to FLASH [M.musculus] {IMAGE:833829}	AA412925.1
2.8	218	ESTs, Highly similar to KIAA0650 protein [H.sapiens] {IMAGE:621629}	AA183687.1
3.2	62	ESTs, Highly similar to major vault protein [R.norvegicus] {IMAGE:6395	AA200827.1
2	67	ESTs, Highly similar to molybdopterin-synthase large subunit [M.musci	AI450527.1
2.2	45	ESTs, Highly similar to myomegalin [R.norvegicus] {IMAGE:315787}	W12392.1
2.2	50	ESTs, Highly similar to NY-REN-2 antigen [H.sapiens] {IMAGE:945689}	AA546195.1
2	47	ESTs, Highly similar to pirin [H.sapiens] {IMAGE:332396}	W08720.1
2.6	142	ESTs, Highly similar to probable nocturnin protein [M.musculus] {IMAGE	AA051133.1
2.6	71	ESTs, Highly similar to putative transcription factor [H.sapiens] {IMAGE	AA467436.1
2	208	ESTs, Highly similar to unknown [H.sapiens] {IMAGE:336564}	AI390954.1

2.3	132	ESTs, Highly similar to unknown [H.sapiens] {IMAGE:492447}	AA097194.1
3.2	87	ESTs, Moderately similar to ACTIN-LIKE PROTEIN 14D [Drosophila mel] AA451385.1	
2	46	ESTs, Moderately similar to GEPHYRIN [Rattus norvegicus] {IMAGE:61} AI893727.1	
2	49	ESTs, Moderately similar to C11D2.4 [C.elegans] {IMAGE:441807} AA014053.1	
2	50	ESTs, Moderately similar to homocysteine respondent protein HCYP2 [I] W20810.1	
2	337	ESTs, Moderately similar to KIAA0651 protein [H.sapiens] {IMAGE:6952} AA220487.1	
2.1	68	ESTs, Moderately similar to probable ligand-binding protein RYD5 [R.n] AA412821.1	
2.1	61	ESTs, Moderately similar to SARCOLIPIN [H.sapiens] {IMAGE:318692} W11587.1	
2.9	61	ESTs, Moderately similar to unknown [H.sapiens] {IMAGE:761038} AA387579.1	
2.1	59	ESTs, Weakly similar to CYP2C39 [M.musculus] {IMAGE:367514} AI427394.1	
2.3	39	ESTs, Weakly similar to cytohesin 2 [M.musculus] {IMAGE:779020} AA413956.1	
2.1	85	ESTs, Weakly similar to CYTOSKELETAL P17 PROTEIN [Dictyostelium] AA017875.1	
2.2	59	ESTs, Weakly similar to DUAL SPECIFICITY PROTEIN PHOSPHATASE 1 AA119029.1	
2.2	623	ESTs, Weakly similar to GARP PROTEIN PRECURSOR [H.sapiens] {IMA AA048874.1} W41083.1	
2.3	43	ESTs, Weakly similar to gob-5 [M.musculus] {IMAGE:351090} W11535.1	
2.1	71	ESTs, Weakly similar to HYPOTHETICAL 11.4 KD PROTEIN C13G6.04 IN W18457.1	
2.2	45	ESTs, Weakly similar to keratin 10, type I, cytoskeletal [H.sapiens] {IMA AA472994.1} W18457.1	
3.1	310	ESTs, Weakly similar to LYMPHOCYTE ANTIGEN LY-6A.2/LY-6E.1 PREC AA472994.1	
3.5	81	ESTs, Weakly similar to lysophospholipase I [M.musculus] {IMAGE:596 AA138854.1} AA209006.1	
2.5	705	ESTs, Weakly similar to mannose-binding lectin associated serine prot W40994.1	
2.3	107	ESTs, Weakly similar to mCAC [M.musculus] {IMAGE:350881} AA541946.1	
2.6	81	ESTs, Weakly similar to ORF YJL004c [S.cerevisiae] {IMAGE:933229} AA518000.1	
2	57	ESTs, Weakly similar to Pro-Pol-dUTPase polyprotein [M.musculus] {IM AA260397.1} W09936.1	
2.1	180	ESTs, Weakly similar to SDP8 [M.musculus] {IMAGE:747180} AA397172.1	
2.2	66	ESTs, Weakly similar to similar to alpha/beta hydrolase fold [C.elegans] AA220505.1	
2.1	76	ESTs, Weakly similar to similar to leucyl-tRNA synthetase [C.elegans] {IMAGE AA080169.1} AA060494.1	
2.3	566	ESTs, Weakly similar to tazarotene-induced gene 2 [H.sapiens] {IMAGE:523380} AA474897.1	
2.3	315	ESTs, Weakly similar to Trif [M.musculus] {IMAGE:523380} AA050167.1	
2.4	298	ESTs, Weakly similar to UBIQUITIN-CONJUGATING ENZYME E2-16 KD AA474897.1	
2.5	591	Extracellular matrix protein 1 {IMAGE:874833} AA050167.1	
2.2	428	Glutamate oxaloacetate transaminase 1, soluble {IMAGE:481381} AA050167.1	

2.2 45 97 Hemochromatosis {IMAGE:313224}

W11889.1

12.5	37	464	Histocompatibility 2, class II antigen A, alpha {IMAGE:747378}	AA272807.1
3.1	62	190	Histocompatibility 2, class II antigen A, beta 1 {IMAGE:618271}	AA175329.1
3.9	59	231	Histocompatibility 2, class II, locus Ma {IMAGE:669933}	AA230649.1
2	329	660	Histocompatibility 2, K region {IMAGE:330336}	W14540.1
2.2	143	314	Histocompatibility 2, Q region locus 7 {IMAGE:596447}	AA122791.1
2.2	311	688	Histocompatibility 2, Q region locus 7 {IMAGE:637849}	AA186012.1
2	560	1133	Histocompatibility 2, T region locus 22 {IMAGE:920425}	AA538511.1
2.1	107	225	Homolog of human MLT2 unidentified gene {IMAGE:750930}	AA289139.1
6.7	36	240	Interferon activated gene 203 {IMAGE:620541}	AA174447.1
4.1	273	1123	Interferon regulatory factor 1 {IMAGE:777393}	AA276697.1
3.4	801	2763	Lymphocyte antigen 6 complex {IMAGE:580715}	AA145865.1
2.7	539	1462	Lymphocyte antigen 6 complex, locus C {IMAGE:425855}	AA000712.1
5.8	37	216	M.musculus mRNA for caspase-11 {IMAGE:618272}	AA175651.1
2.5	534	1329	M.musculus mRNA for mTGF protein {IMAGE:474339}	AA038657.1
3.3	322	1066	M.musculus mRNA for mTGF protein {IMAGE:722623}	AA260654.1
2.3	61	140	M.musculus spasmolytic polypeptide (SP) mRNA {IMAGE:314078}	W09829.1
2.1	47	99	Metallothionein 2 {IMAGE:427339}	AA003303.1
49.5	62	3070	Monokine induced by gamma interferon {IMAGE:874758}	AA472492.1
3	557	1655	Mouse chromatin nonhistone high mobility group protein (HGM-I(Y), co	AA538243.1
2.3	94	216	Mus musculus A10 mRNA, partial cds {IMAGE:333376}	W15888.1
2.2	401	902	Mus musculus A10 mRNA, partial cds {IMAGE:385441}	W61383.1
2	72	145	Mus musculus Ankhzn mRNA, complete cds {IMAGE:894596}	AA517717.1
2	349	710	Mus musculus asparagine synthetase mRNA, complete cds {IMAGE:33:	W29492.1

2	147	287	<i>Mus musculus</i> BS4 peptide mRNA, complete cds {IMAGE:459091}	AA024028.1
2	162	317	<i>Mus musculus</i> CIPER protein (Ciper) mRNA, complete cds {IMAGE:355: W47752.1}	W47752.1
2.1	64	137	<i>Mus musculus</i> classII MHC transactivator CIITA mRNA, complete cds {IMAGE:651: AA178010.1}	AA178010.1
2.1	400	836	<i>Mus musculus</i> G-protein-like LRG-47 mRNA, complete cds {IMAGE:651: AA214784.1}	AA214784.1
2.1	36	74	<i>Mus musculus</i> IER5 (Ier5) mRNA, complete cds {IMAGE:643745}	AA203798.1
2.4	57	136	<i>Mus musculus</i> interferon regulatory factor 5 (mirf5) mRNA, complete cds {IMAGE:643745}	AA185025.1
2.53.7	29	1558	<i>Mus musculus</i> interferon-induced guanylate binding protein GBP-2 (Gbp)	AA153021.1
2.4	44	107	<i>Mus musculus</i> intersectin-EH binding protein Ibp2 mRNA, partial cds {IMAGE:621346}	AA413761.1
2.1	141	291	<i>Mus musculus</i> ion channel homolog RIC mRNA, complete cds {IMAGE:621346}	W98807.1
2.2	59	127	<i>Mus musculus</i> major histocompatibility complex region NG27, NG28, R	W13905.1
2.5	36	90	<i>Mus musculus</i> mRNA for DNA helicase Q1, complete cds {IMAGE:3304: AI390823.1}	AI390823.1
6.4	39	249	<i>Mus musculus</i> mRNA for GTP1 protein {IMAGE:621346}	AA178613.1
20.2	35	708	<i>Mus musculus</i> mRNA for IIGP protein {IMAGE:747208}	AA260490.1
20.9	50	1044	<i>Mus musculus</i> mRNA for IIGP protein {IMAGE:764385}	AA277451.1
2	42	82	<i>Mus musculus</i> mRNA for MD-1, complete cds {IMAGE:402964}	W80147.1
2	49	99	<i>Mus musculus</i> peroxisomal biogenesis factor (Pex11a) mRNA, complete	AA237383.1
10	81	809	<i>Mus musculus</i> putative purine nucleotide binding protein mRNA, complete	AA240404.1
2.3	128	288	<i>Mus musculus</i> secreted carbonic anhydrase isozyme VI precursor, mRNA	AI327498.1
2.5	535	1362	Myelocytomatosis oncogene {IMAGE:441346}	AA009268.1
2.7	234	625	Myeloid differentiation primary response gene 118 {IMAGE:386412}	W64388.1
3.2	520	1650	MYELOID DIFFERENTIATION PRIMARY RESPONSE PROTEIN MYD116 {IMAGE:386412}	AA050417.1
2	352	719	NAD-DEPENDENT METHYLENETETRAHYDROFOLATE DEHYDROGENA	W84014.1
3.2	507	1629	Protease (prosome, macropain) 28 subunit, alpha {IMAGE:678768}	AA239485.1
2	83	162	Protein tyrosine phosphatase, non-receptor type 2 {IMAGE:437517}	AA003599.1
2.2	48	105	Public domain EST {IMAGE:314509}	W10776.1
2.2	49	110	Public domain EST {IMAGE:314524}	W10780.1
2.3	53	122	Public domain EST {IMAGE:330298}	W13940.1
2	53	106	Public domain EST {IMAGE:350327}	W40724.1
2.4	82	194	Public domain EST {IMAGE:353202}	W41810.1

109	Public domain EST {IMAGE:367869}
53	Public domain EST {IMAGE:372366}
42	Public domain EST {IMAGE:374745}
63	Public domain EST {IMAGE:374745}
2	Public domain EST {IMAGE:390150}
39	Public domain EST {IMAGE:403961}
2.1	Public domain EST {IMAGE:439362}
41	Public domain EST {IMAGE:442201}
2.2	Public domain EST {IMAGE:456831}
2	Public domain EST {IMAGE:459599}
131	Public domain EST {IMAGE:463183}
2.1	Public domain EST {IMAGE:465170}
2.2	Public domain EST {IMAGE:466376}
2	Public domain EST {IMAGE:478309}
58	Public domain EST {IMAGE:479480}
2.3	Public domain EST {IMAGE:482196}
2	Public domain EST {IMAGE:482817}
51	Public domain EST {IMAGE:574439}
2.3	Public domain EST {IMAGE:596231}
50	Public domain EST {IMAGE:597020}
3.3	Public domain EST {IMAGE:617529}
4.3	Public domain EST {IMAGE:618199}
3.3	Public domain EST {IMAGE:619596}
4.7	Public domain EST {IMAGE:620030}
108	Public domain EST {IMAGE:621742}
4.7	Public domain EST {IMAGE:636640}
13.8	Public domain EST {IMAGE:652811}
4.6	Public domain EST {IMAGE:653161}
108	Public domain EST {IMAGE:733734}
4.6	Public domain EST {IMAGE:760824}
77	Public domain EST {IMAGE:760917}
66	Public domain EST {IMAGE:761009}
62	Public domain EST {IMAGE:774996}

4.6	50	231	Public domain EST	{IMAGE:775154}	AI604749.1
2.2	44	97	Public domain EST	{IMAGE:778923}	AA414282.1
2	44	86	Public domain EST	{IMAGE:779453}	AA414212.1
2.2	67	146	Public domain EST	{IMAGE:805306}	AA473329.1
2.5	71	180	Public domain EST	{IMAGE:948845}	AA544859.1
2.7	86	229	Recombining binding protein suppressor of hairless (Drosophila)	{IMAGE:W97158.1}	W97158.1
2.3	39	88	SAPK/Erk/kinase 1	{IMAGE:778459}	AA413508.1
7.4	116	859	Signal transducer and activator of transcription 1	{IMAGE:749544}	AA388678.1
2.2	52	113	Thymus cell antigen 1, theta	{IMAGE:316409}	W13151.1
15.7	40	629	Transporter 1, ABC (ATP binding cassette)	{IMAGE:576775}	AA123837.1
2.3	45	104	Transthyretin	{IMAGE:335319}	W17647.1
7.7	198	1527	TRYPTOPHANYL-TRNA SYNTHETASE	{IMAGE:367765}	W53959.1
2.6	401	1028	Tumor necrosis factor induced protein 2	{IMAGE:949472}	AA537672.1
2.2	80	175	Zinc finger protein 35	{IMAGE:422121}	AI893996.1
2.3	82	190	Zinc transporter 4	{IMAGE:643619}	AA203866.1

72 genes up >= 2-fold 022PAV3Q (Exp M subm 2/17/00)

Fold	Cont	Tg	Gene name	Acc#
2.7	527	1434	apoptosis Inhibitor 1 {IMAGE:640046}	AA197349.1
2.1	279	581	B-cell leukemia/lymphoma 10 {IMAGE:355209}	W47752.1
2.5	2462	6113	beta-2 microglobulin {IMAGE:572542}	AA109951.1
2.1	609	1263	capping protein alpha 1 {IMAGE:779754}	AA414612.1
3.1	125	388	caspase 11 {IMAGE:618272}	AA175651.1
2.1	1054	2215	cell division cycle 42 homolog (S. cerevisiae) {IMAGE:721518}	AA266975.1
3.4	321	1087	ESTs {IMAGE:523814}	AA067146.1
3.4	190	639	ESTs {IMAGE:583566}	AA143986.1
9.6	369	3541	ESTs {IMAGE:595951}	AA140542.1
2	150	298	ESTs {IMAGE:596556}	AA146363.1
3.9	165	647	ESTs {IMAGE:596863}	AA145136.1
2.5	108	265	ESTs {IMAGE:597031}	AA145466.1
5.6	148	827	ESTs {IMAGE:619723}	AA172624.1
3.5	125	441	ESTs {IMAGE:636640}	AA204588.1
2	556	1133	ESTs {IMAGE:636827}	AA183302.1
5.7	366	2072	ESTs {IMAGE:678854}	AA250462.1
3.1	306	954	ESTs {IMAGE:751826}	AA396152.1
3.3	162	542	ESTs {IMAGE:777379}	AA277040.1
2.5	368	928	ESTs {IMAGE:876257}	AA476183.1
2	237	484	ESTs, Highly similar to HYPOTHETICAL 103.5 KD PROTEIN IN RAD26-GI	AI639807.1
2.2	224	502	ESTs, Highly similar to INTERFERON-ACTIVATABLE PROTEIN 204 [Mus	AI591873.1
2.8	866	2458	ESTs, Highly similar to INTERFERON-INDUCIBLE PROTEIN [Rattus norv	AA048837.1
2.1	243	511	ESTs, Highly similar to AE-1 binding protein AEBP2 [R.norvegicus] {IMAGE:622724}	AA416308.1
2.2	556	1207	ESTs, Highly similar to Bdelight protein [R.norvegicus] {IMAGE:622724}	AA177555.1
2.5	426	1070	ESTs, Highly similar to probable nocturnin protein [M.musculus] {IMAGE:AA051133.1}	
7.2	293	2099	ESTs, Highly similar to purine nucleotide binding protein [M.musculus] {IMAGE:AA240404.1}	
2.6	423	1108	ESTs, Highly similar to spilloosomal protein SAP 155 [H.sapiens] {IMAGE:AA200809.1}	

3.6	124	451	ESTs, Moderately similar to ACTIN-LIKE PROTEIN 14D [Drosophila mela	AA451385.1
3.3	663	2170	ESTs, Moderately similar to CHLORINE CHANNEL PROTEIN P64 [Bos tauri]	AA466508.1
2	197	391	ESTs, Moderately similar to CHLORINE CHANNEL PROTEIN P64 [Bos tauri]	AA537234.1
2.4	188	442	ESTs, Moderately similar to PLASMA-CELL MEMBRANE GLYCOPROTEI	AA016397.1
2.2	680	1508	ESTs, Moderately similar to PRE-MRNA SPLICING FACTOR SRP20 [Hom	W83358.1
2.2	330	740	ESTs, Weakly similar to CCAAT-BINDING TRANSCRIPTION FACTOR SUF	AA413894.1
2.9	785	2288	ESTs, Weakly similar to LYMPHOCYTE ANTIGEN LY-6A/2/LY-6E.1 PREC	AA472994.1
2.5	160	403	ESTs, Weakly similar to lysophospholipase I [M.musculus]	{IMAGE:5964; AA153793.1}
2.8	1172	3251	ESTs, Weakly similar to mannose-binding lectin associated serine protease	AA209006.1
2.8	625	1738	ESTs, Weakly similar to SDP8 [M.musculus]	{IMAGE:747180}
2.1	413	871	ESTs, Weakly similar to Trif [M.musculus]	{IMAGE:523380}
3.3	214	706	expressed sequence tag mouse EST 25	{IMAGE:638284}
2.1	713	1511	interferon activated gene 203	{IMAGE:618572}
2.7	255	697	interferon activated gene 203	{IMAGE:620541}
4.8	1249	5945	interferon inducible protein 1	{IMAGE:651927}
9.6	411	3937	interferon regulatory factor 1	{IMAGE:777393}
2.8	2057	5726	lymphocyte antigen 6 complex	{IMAGE:580715}
2.5	792	1977	lymphocyte antigen 6 complex, locus C	{IMAGE:425855}
4.4	225	980	monokine induced by gamma interferon	{IMAGE:874758}
21	204	4277	Mus musculus interferon-induced guanylate binding protein GBP-2 (Gbp	AA153021.1
2.8	279	777	Mus musculus mRNA for GTP1 protein	{IMAGE:621346}
2	1359	2725	Mus musculus mRNA for heterogeneous nuclear ribonucleoprotein H	{IM AA003085.1}
4.4	125	549	Mus musculus mRNA for IIGP protein	{IMAGE:747208}
5.6	229	1293	Mus musculus mRNA for IIGP protein	{IMAGE:764385}
2	268	546	myeloid differentiation primary response gene 88	{IMAGE:572317}
4.8	228	1096	nuclear factor of kappa light chain gene enhancer in B-cells Inhibitor, alp	AA204405.1
2	606	1236	protein kinase C, delta	{IMAGE:777583}
3.6	184	664	Public domain EST	{IMAGE:482196}
2.9	132	379	Public domain EST	{IMAGE:596231}

5.8	105	614	Public domain EST {IMAGE:597020}	
7.1	221	1571	Public domain EST {IMAGE:617529}	AA138757.1
2	122	247	Public domain EST {IMAGE:619686}	AA174721.1
3.4	138	471	Public domain EST {IMAGE:653161}	AA175973.1
2	215	425	Public domain EST {IMAGE:669831}	AA1390695.1
2.3	411	935	Public domain EST {IMAGE:733734}	AA253750.1
2.1	383	801	Public domain EST {IMAGE:850178}	AA272876.1
2.7	1687	4542	Public domain EST {IMAGE:874030}	AA451403.1
2.6	579	1531	purine-nucleoside phosphorylase {IMAGE:607469}	AA472200.1
7.8	222	1740	signal transducer and activator of transcription 1 {IMAGE:749544}	AA172867.1
3	617	1833	signal transducer and activator of transcription 3 {IMAGE:440571}	AA388678.1
3	666	1990	small inducible cytokine subfamily D, 1 {IMAGE:761708}	AA014643.1
4.4	138	613	transporter 1, ABC (ATP binding cassette) {IMAGE:576775}	AA272232.1
2.2	329	719	tryptophanyl-tRNA synthetase {IMAGE:367765}	AA123837.1
2	357	717	Y box protein 3 {IMAGE:421546}	W53959.1
2.4	944	2275	zinc finger protein 36 {IMAGE:641523}	W97158.1
				AA2000282.1